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Title: Towards establishing consistency in triage in a tertiary specialty

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Running Title: Multicentre Review of Triage in Clinical Genetics

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Keywords: Triage; Prioritisation; referrals; clinical genetics

Abstract

Clinical Genetics services provide a diagnostic, counselling and genetic testing service for children and adults affected by, or at risk of, a genetic condition, most of which are rare, or genetically heterogeneous. Appropriate triage of referrals is crucial to ensure the most urgent referrals are seen as quickly as possible, without negatively impacting the waiting times of less urgent cases. We aimed to examine triage practice in 6 Clinical Genetic centres across the UK and Ireland. Thirteen simulated referrals were drafted based on common referrals to Clinical Genetics. Copies of each referral were forwarded to each centre, where 10 nominated clinicians were asked to triage each referral. Triage referrals were returned to the coordinating author for analysis. An electronic questionnaire was contemporaneously completed by clinical leads in each unit to gather local demographic details and local operating procedures relevant to triage. Widespread inconsistencies were noted both within and between units, with respect to acceptance of referrals to services, prioritisation, and designated clinic type. Referral rates, staffing levels, and waiting lists varied widely between units. Inconsistencies observed between units are likely influenced by a number of factors including; staffing levels, referral rates, and average family size. Inconsistency within units likely reflects the complex nature of many Clinical Genetic referrals and triage guidelines should help improve decision making in this setting.

21 Introduction

22 Clinical Genetics services provide a diagnostic, counselling and genetic testing service
23 for children and adults affected by, or at risk of, a genetic condition[1]. Referrals come
24 from almost all specialties, from primary, secondary and tertiary centres[2]. The
25 geographical catchment area, and indications for referral (from neonatal to adult;
26 dysmorphology, and referrals from all subspecialties) covered by Clinical Genetics
27 centres is wide.

28 Increasingly broad genetic testing has led to discovery of novel disease genes, and new
29 genotype-phenotype associations (figure 1)[3]. This has positively impacted diagnostic
30 yield in patients with disorders related to previously undefined genetic aetiology (e.g.
31 epilepsy, sudden adult death); but has also led to increased detection of variants of
32 uncertain significance[4, 5], and of variants in genes not previously known to be
33 associated with a particular phenotype (“genes of uncertain significance”)[6]. Such
34 variants generate massive clinical workload, and often require reviewing multiple
35 family members to facilitate segregation analysis; or may require multiple patient
36 encounters to facilitate collection of different sample types for functional studies (e.g.
37 skin or muscle biopsy, biochemical testing). A single referral may therefore generate
38 many days-weeks of clinical work. Furthermore, absolute numbers of referrals may be
39 a poor reflection of the workload of a unit, depending on the complexity of the case-
40 mix [7]. Benign, likely benign or uncertain variants are frequently picked up by array
41 CGH[8], a test routinely used by general paediatricians. As non-Geneticists grapple
42 with increasingly complex genetic test reports they request advice to help interpret
43 the report; while the actual presenting complaint in the patient may have been

considered too trivial to refer in the past. Previous audits suggest referrals to explain normal benign or likely benign human variation account for 10% of general referrals [9].

The specialty mainly receives non-urgent out-patient referrals, however, pre-natal referrals, or referrals for patients approaching end of life require prompt assessment. Demand for urgent access to genetic testing is growing where results might influence management. Increasingly, targeted therapies are being licenced for use in patients with germline or somatic genetic variation, particularly in treatment of cancer (e.g. PARP inhibitors, ATR inhibitors, small molecule kinase inhibitors)[10, 11]. Public and media awareness has also driven demand, both those affected or at risk of a familial genetic disorder[12]. Increasing cost-efficiency of testing has led to an interest in population-based screening for genetic disorders[13-15], and has driven direct-to-consumer testing, with predicted market value of up to \$310 million by 2022 [16, 17]. Consequently, this puts increasing stress on under-resourced genetic services.

Genetic counsellors are highly skilled clinical professionals, usually from scientific or nursing backgrounds, with specialist training in genetic counselling[18]. Not all countries employ genetic counsellors, but they form a core part of the Clinical Genetics teams in the UK and Ireland[19]. In most genetic centres in the UK and Ireland, Consultant Geneticists review undiagnosed or complex patients, while genetic counsellors review patients at risk of a known familial genetic disorder, to offer pre-symptomatic predictive testing. Some centres utilise a co-counselling approach involving both types of professional[19], while in other centres, patients have an initial

“pre-clinic” with a genetic counsellor, followed thereafter by consultant-led interaction. It is well-recognised that there is a significant shortage of both Genetic Counsellors and Consultant Clinical Geneticists internationally, particularly in Ireland and England [20-22]. Appropriate triage of referrals is a critical factor in trying to address demands on the service in the face of limited resources; to ensure the most urgent referrals are seen as quickly as possible, without negatively impacting the waiting times of less urgent cases. To ensure optimal provision of services, the Clinical Genetics Society has considered a number of common referrals that do not need face-to-face consultation in a Clinical Genetics Centre[23]. Centres have also adopted local policies to reject referrals pertaining to conditions where specialist clinics exist in the region [24]. In Centre 1, for example, all referrals related to patients with inherited cardiac pathologies are deferred to the Cardiology service. In Centre 3, referrals related to common paediatric conditions such as Down Syndrome or Spina Bifida are managed by letter to the patient, without offering patient a formal consultation. This may partly explain inter-departmental differences.

However, as referrals may pertain to any one of thousands of different rare disorders, standardisation of referrals is very difficult. We aimed to review the practice of triage in Clinical Genetics centres in the UK and Ireland using high-fidelity simulated referrals.

Methods

A consultant geneticist in each centre was identified and asked to co-ordinate the study locally. Participants were asked to complete a short questionnaire to establish local demographics and local practice at their respective centre. Data was collected

with respect to factors that could potentially influence triage practice, including staffing level, waiting lists, catchment area and population size, clinician responsible for triage, and number of referrals per year.

Thirteen simulated referrals were designed (by TMcV and SAL). Ten were based on genuine referrals, with patient, referring Doctor and hospital identifiers removed, and details changed slightly so as to maintain confidentiality in line with European General Data Protection Regulation legislation. The remaining 3 (referral no. 4, 7 & 13) were composed by the authors based on common referrals to a Clinical Genetics service. All were printed on headed notepaper of a fictitious hospital (Supplementary Figure 1), and 10 hard copies were posted to each centre. This was to endeavour to create high-fidelity simulated referrals on the expectation that the research triage would be a true reflection of genuine triage [25, 26]. The nature of the 13 referrals can be seen in table

1. Participants were told these were simulated referrals. We deliberately mis-spelt certain words, and inserted information regarding a patient's pregnancy in the middle of a referral rather than placing emphasis on the urgency of the referral, reflecting frequent errors in referrals from practitioners unfamiliar with genetic conditions and implications of such disorders for progeny of affected individuals, increasing fidelity of the simulation.

Participants were asked to triage each referral by type of appointment; urgency; designated clinician, etc, using a standardised triage stamp (figure 2). Completed triage forms were posted back to the lead author in the coordinating centre. Data were tabulated and analysed using SPSS v23.

Results

Participants

In total, 53 clinicians from 6 centres participated in the simulated triage exercise. Participants included 27 consultants (51%), 19 Genetic Counsellors (36%) and 7 (13%) Specialist Registrars (table 2). All participants from centre 5 were consultants as local practice dictates that only consultants perform triage. In Centres 4 and 6, certain consultants perform triage for both General and Cancer cases, while others triage only one category or the other. Depending on their local practice, some clinicians declined to triage certain simulated referrals.

Significant variability in the process of triage was noted across the six centres. In three centres (Centres 1, 4 and 5), triage of general referrals was undertaken by consultants only, and in the other three centres, by consultants and GCs. Triage of cancer referrals was, conversely, done by GCs only in three centres (Centres 1,2 and 4), and by consultant only in Centres 5 and 6. All centres accept referrals by letter. Five centres accept electronic referrals, and four accept referral by fax (table 3).

Between centres, there was variability in the number of referrals per 1000 of population per annum (0.84-3/1000), and the number of referrals per consultant and per staff member, which could not be explained by average family size. Centre 3 and Centre 1 had almost equivalent numbers of referrals despite >2.5-fold difference in the size of population.

There were clear discrepancies in staffing numbers with Centre 2 being relatively well staffed and Centre 3 being very poorly staffed, with respect to both consultant and GC workforce. The ratio of referrals/staff member was lowest in Centre 4, and highest in Centre 1. The proportion of referrals managed without a face-to-face appointment was highest in Centre 4 and lowest in Centre 1 (8-38).

Acceptance of referrals to service

Considering all clinicians, widespread variability in triage was noted (Figure 3 (3a)). Only 3 (23%) of the referrals had >80% consensus about whether the referral should be accepted for a consultation. There was complete or almost complete consistency (>80% consensus) with the triage decision for five referrals (referrals 1,3, 5, 6 and 10) amongst consultants (Figure 3 (3b)), and consensus of 60-80% for three others (referrals 9, 12 and 13).

Significant inconsistency was noted for the other referrals, with some consultants offering a face to face appointment, and others managing the same referrals by providing an information letter or telephone consultation to the patient. Other clinicians elected to reject the referral and provide referrer with information about onward management of the patient, without direct patient contact.

When triage performed by Genetic counsellors was considered, only two referrals (referral 1 and 10) had >80% consensus regarding type of consultation offered. Referral 3,4 and 6 showed 60-80% consensus (Figure 3 (3c)). Consensus between and within centres are demonstrated in supplementary figures 2-12.

Prioritisation of Referrals

Of those referrals offered face to face appointments, significant variability in priority and designated clinician was also noted (supplementary table 1). In a significant number of cases, clinicians did not specify priority/designated clinician (excluded).

The referral with the most agreement between clinicians was a simulated urgent referral of a pregnant woman with a family history of Duchenne muscular dystrophy. Forty-eight clinicians triaged this referral, and all accepted the referral to service. 47/48 specified the priority of the referral as priority (1 did not specify). There was inconsistency in determining designated clinician, with 26 (57%) triaging the case for GC appointment, and 20 (43%) for consultant (2 did not specify).

Consistency within Centres (Supplementary Table 1)

Where appointments were offered, 100% consistency was noted for prioritisation of five referrals, including referral 10 as priority; as well as referrals 8, 9, 11¹ and 13 which were deemed routine by all participants. Certain centres were more consistent than others with respect to prioritisation of those referrals for which face-to-face appointments were offered. Clinicians in Centre 2 agreed on priority of an additional two referrals of those 8

¹ In one centre, referrals pertaining to Hereditary Haemochromatosis are deferred to Gastroenterology so are not offered appointments in the Clinical Genetics unit

169 offered face to face appointments in that centre (25%), Centre 3 2/5 (40%), Centre 5, Centre
170 6 and Centre 1 another 4/7 (57%), and Centre 4 5/8 (63%).

171 With respect to designated clinicians, inconsistency across each referral was noted. In
172 Centre 3, of 8 referrals offered appointments, there was agreement between participants
173 there for designated clinician in 5 (63%) appointments. In Centre 2 and Centre 4, all referrals
174 would be offered appointments by at least one clinician², but there was agreement in these
175 centres with respect to designated clinician in only 2 (15%) cases.

² Referral 7 was offered an appointment by only one clinician in Centre 2, and by two clinicians in Centre 4

176

177 Discussion

178 Clinical triage is an important step in all specialties, aiming to ensure prioritisation of
179 referrals and maintain equity of access. Our study has shown widespread inconsistencies in
180 managing common referral scenarios both within and between six Clinical Genetic units in
181 the Republic of Ireland and the United Kingdom. Inconsistencies were noted with respect to
182 acceptance of referrals to service, prioritisation of referrals, and type of clinic to which
183 referral was assigned

184 Discrepancy between centres with respect to type of consultation offered to patients may
185 be attributed to hospital management systems; in the Republic of Ireland; referrals that
186 were not offered a face-to-face consultation were deemed rejected, despite providing the
187 patient with information directly, whereas similar practices in centres in UK system were
188 acknowledged, and remunerated, as clinical activity. However, this does not explain the
189 differences between clinicians within centres. Differences in priority assigned to cases may
190 be influenced by waiting lists and staffing, which vary between centres. It is possible that
191 decision-making with respect to assignment of cases to Consultant or Genetic Counsellor
192 may be influenced by the level of expertise of staff within the unit.

193 Traditionally, research on triage has concentrated on pre-hospital, trauma, acute or
194 emergency care settings[27-31]. Assessment of triage in tertiary referrals specialties has also
195 concentrated on optimising management in the acute scenario[32]. Appropriate triage in
196 tertiary referral setting is important to ensure equity of care, timely access based on need
197 and an ability to manage waiting lists in accordance with staffing levels[33-36]. Prioritisation
198 of the most urgent referrals is critical when waiting lists deteriorate and timely access to

199 care is at risk[37-39]. Each speciality will have specific drivers that influence the ebb and
200 flow of referrals. Triage decision making in Clinical Genetics is driven by many factors
201 related to the centre in question (e.g. staffing levels, skill mix, waiting list times, population
202 demographics), the patient to which referral pertains (e.g. pregnant patient, patient
203 approaching end of life, patient age, patient at risk of inheriting familial variant), or nature
204 of the referral itself (request for genetic information to determine treatment, advice to
205 interpret genetic test results, adequacy of information on referral letter).

206 Factors known to influence referral rates include education of referrers, the genetics
207 workforce, and logistic factors[40, 41]. We noted regional differences in referral rates/1000
208 population, which have not previously been described. A number of factors may account for
209 these apparent differences. Genetic disorders may be more prevalent in countries where
210 there are endogamous populations (e.g. Irish Travellers), with associated founder mutations
211 and disorders[42]; and among populations where first-cousin marriage is permitted, with
212 associated increased incidence of recessive disorders. Birth rates in the Republic of Ireland
213 (13.5 per 1000) and Northern Ireland (13.1/1000) are higher than the reported 11.8/1000 in
214 England, Scotland and Wales and these together with the current lack of availability of
215 termination of pregnancy on the island of Ireland result in more urgent liveborn referrals
216 which may impact regional differences in referral rates.

217 The March of Dimes describes that a fundamental role of a Clinical Genetic service is
218 prevention [43]. One component of this is to offer cascade genetic testing to at-risk relatives
219 of patients with confirmed genetic disorders. Many referrals may therefore be generated by
220 a single family once a pathogenic genetic variant is identified. Cascade screening is
221 particularly burdensome in countries with large family sizes. In Ireland, the average size of

222 an extended 3 generation family [including siblings of grandparents and their offspring] is >3
223 times (64 vs 19) that of average families in England/Wales and Scotland (figure 4) [44, 45].
224 Unsurprisingly, cascade screening for common dominant genetic disorders accounts for 12%
225 of general referrals in Ireland. Regional differences in referral rates may be further
226 explained by differences in management of such referrals. In some centres, at risk relatives
227 may self-refer by telephone or email. Furthermore, other relatives may opt to attend the
228 appointment offered to one individual in the family to “skip the queue”. Generally, these
229 patients are facilitated, counselled and treated, but may not be recorded as a “referral”.
230 Other centres require formal referrals from GPs or secondary care to facilitate review of
231 relatives for cascade testing.

232

233 Population demographics and local policies cannot completely account for the lack of
234 consistency within units. In certain units, genetic counsellors perform “pre-clinic” for
235 consultants; and this may explain the variability in assignment of designated consultant.
236 Some participants may have selected “genetic counsellor” based on the first appointment to
237 which referral would be assigned, while others may have selected “consultant” as referral
238 would ultimately end up in a consultant clinic. In centres where genetic counsellor staffing
239 levels are sub-optimal, pre-clinics are not possible.

240 In most specialties, priority is defined by urgency of the referral, which may not be
241 appropriate in specialties like Clinical Genetics, where most referrals are non-urgent[46].
242 Defining priority by urgency may therefore disadvantage the majority of patients referred to
243 Clinical Genetics[39], putting routine waiting lists under strain. There are no current
244 guidelines one can use to determine priority of referral, although a shared set of

prioritisation criteria have been proposed – including the clinical and non-clinical benefits to patient and family; risk, progression, and severity of disease, and cost, and infrastructure for testing [47, 48].

In our study, it is likely that other factors, such as local waiting lists, availability of regional specialist clinics, or human subjectivity may explain the inconsistencies we have observed within each centre. Certain specific situations (e.g. if genetic diagnosis required prior to undergoing surgery, starting new treatment, accessing services etc.) may mean that cases that might otherwise be rejected or managed by letter will be offered face-to-face appointments. All centres involved in this study are training centres for Clinical Geneticists, and common conditions that might otherwise be deflected to another specialty might be accepted to fulfil curricular requirements.

[Limitations of this study](#)

Each individual centre faces different pressures with respect to staffing and waiting lists, which will, in turn, impact triage practice. It is possible that the process might differ in each centre when dealing with real referrals, all participants knew this was a research study and that referrals were high-fidelity simulations; participants may therefore have been more casual in their answers. We did not collect data specifically with respect to waiting times for routine or priority appointments. We note the NHS guideline of maximum 18 weeks, but appreciate that many centres in the UK struggle to avoid breaching this timeline. As a direct consequence of poor staffing levels, in the Republic of Ireland, the waiting times for priority appointments are in the order of 12-14 months, and for routine, 18-24 months. Attempts to recruit and retain trainees and genetic counsellors; and upskill non-genetic specialist is a continuing challenge.

Conclusion

The consensus in triage established in this paper should form the basis for guidelines to help an equitable consistent approach to these 13 common referrals. Individual centres will need to establish more standardised local policies in the context of their own staffing levels and availability of regional specialist clinics, but national/international guidelines are required to ensure equity in the triage process. We are mindful that we have examined the process with 13 common referrals; ensuring consistency is likely to be even more challenging when addressing the complex referrals received by all clinical genetics service. We would suggest that this issue should be considered in a European context; possibly by convening a workshop at the European Society of Human Genetics annual meeting.

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